

X-13268

SEQUENCE LISTING~

<110> Witcher, Derrick
 Tian, Yu
 Atkinson, Paul

<120> FLINT Analog Compounds and Formulations Thereof

<130> X-13268

<140>

<141>

<160> 3

<170> PatentIn Ver. 2.0

<210> 1

<211> 271

<212> PRT

<213> Homo sapiens

<400> 1

Val	Ala	Glu	Thr	Pro	Thr	Tyr	Pro	Trp	Arg	Asp	Ala	Glu	Thr	Gly	Glu
1				5					10					15	
Arg	Leu	Val	Cys	Ala	Gln	Cys	Pro	Pro	Gly	Thr	Phe	Val	Gln	Arg	Pro
			20					25					30		
Cys	Arg	Arg	Asp	Ser	Pro	Thr	Thr	Cys	Gly	Pro	Cys	Pro	Pro	Arg	His
		35					40					45			
Tyr	Thr	Gln	Phe	Trp	Asn	Tyr	Leu	Glu	Arg	Cys	Arg	Tyr	Cys	Asn	Val
	50					55					60				
Leu	Cys	Gly	Glu	Arg	Glu	Glu	Glu	Ala	Arg	Ala	Cys	His	Ala	Thr	His
65					70				75						80
Asn	Arg	Ala	Cys	Arg	Cys	Arg	Thr	Gly	Phe	Phe	Ala	His	Ala	Gly	Phe
				85					90					95	
Cys	Leu	Glu	His	Ala	Ser	Cys	Pro	Pro	Gly	Ala	Gly	Val	Ile	Ala	Pro
			100					105					110		
Gly	Thr	Pro	Ser	Gln	Asn	Thr	Gln	Cys	Gln	Pro	Cys	Pro	Pro	Gly	Thr
		115					120					125			
Phe	Ser	Ala	Ser	Ser	Ser	Ser	Ser	Glu	Gln	Cys	Gln	Pro	His	Arg	Asn
	130						135				140				
Cys	Thr	Ala	Leu	Gly	Leu	Ala	Leu	Asn	Val	Pro	Gly	Ser	Ser	Ser	His
145					150					155					160
Asp	Thr	Leu	Cys	Thr	Ser	Cys	Thr	Gly	Phe	Pro	Leu	Ser	Thr	Arg	Val
				165					170					175	
Pro	Gly	Ala	Glu	Glu	Cys	Glu	Arg	Ala	Val	Ile	Asp	Phe	Val	Ala	Phe
			180					185					190		

X-13268

Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu
195 200 205
Ala Pro Glu Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu
210 215 220
Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp
225 230 235 240
Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met
245 250 255
Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His
260 265 270

<210> 2
<211> 300
<212> PRT
<213> Homo sapiens

<400> 3
Met Arg Ala Leu Glu Gly Pro Gly Leu Ser Leu Leu Cys Leu Val Leu
1 5 10 15
Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly Val Ala Glu
20 25 30
Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu Arg Leu Val
35 40 45
Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg
50 55 60
Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln
65 70 75 80
Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val Leu Cys Gly
85 90 95
Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His Asn Arg Ala
100 105 110
Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu
115 120 125
His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro
130 135 140
Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala
145 150 155 160
Ser Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala
165 170 175
Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu
180 185 190
Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala

X-13268

195	200	205
Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile 210 215 220		
Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu 225 230 235 240		
Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys 245 250 255		
Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu 260 265 270		
Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu 275 280 285		
Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 290 295 300		

<210> 3
<211> 936
<212> DNA
<213> Homo sapiens

<220>
<221> CDS
<222> (25)..(924)

<400> 3 gctctccctg ctccagcaag gacc atg agg gcg ctg gag ggg cca ggc ctg Met Arg Ala Leu Glu Gly Pro Gly Leu 1 5	51
tgc ctg ctg tgc ctg gtg ttg gcg ctg cct gcc ctg ctg ccg gtg ccg Ser Leu Leu Cys Leu Val Leu Ala Leu Pro Ala Leu Leu Pro Val Pro 10 15 20 25	99
gct gta cgc gga gtg gca gaa aca ccc acc tac ccc tgg cgg gac gca Ala Val Arg Gly Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala 30 35 40	147
gag aca ggg gag cgg ctg gtg tgc gcc cag tgc ccc cca ggc acc ttt Glu Thr Gly Glu Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe 45 50 55	195
gtg cag cgg ccg tgc cgc cga gac agc ccc acg acg tgt ggc ccg tgt Val Gln Arg Pro Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys 60 65 70	243
cca ccg cgc cac tac acg cag ttc tgg aac tac ctg gag cgc tgc cgc Pro Pro Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg 75 80 85	291
tac tgc aac gtc ctc tgc ggg gag cgt gag gag gag gca cgg gct tgc Tyr Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys 90 95 100 105	339

X-13268

cac gcc acc cac aac cgt gcc tgc cgc tgc cgc acc ggc ttc ttc gcg	387
His Ala Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala	
110 115 120	
cac gct ggt ttc tgc ttg gag cac gca tgc tgt cca cct ggt gcc ggc	435
His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly	
125 130 135	
gtg att gcc ccg ggc acc ccc agc cag aac acg cag tgc cag ccg tgc	483
Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys	
140 145 150	
ccc cca ggc acc ttc tca gcc agc agc tcc agc tca gag cag tgc cag	531
Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser Ser Glu Gln Cys Gln	
155 160 165	
ccc cac cgc aac tgc acg gcc ctg ggc ctg gcc ctc att gtg cca ggc	579
Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala Leu Ile Val Pro Gly	
170 175 180 185	
tct tcc tcc cat gac acc ctg tgc acc agc tgc act ggc ttc ccc ctc	627
Ser Ser Ser His Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu	
190 195 200	
agc acc agg gta cca gga gct gag gag tgt gag cgt gcc gtc atc gac	675
Ser Thr Arg Val Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp	
205 210 215	
ttt gtg gct ttc cag gac atc tcc atc aag agg ctg cag cgg ctg ctg	723
Phe Val Ala Phe Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu	
220 225 230	
cag gcc ctc gag gcc ccg gag ggc tgg gct ccg aca cca agg gcg ggc	771
Gln Ala Leu Glu Ala Pro Glu Gly Trp Ala Pro Thr Pro Arg Ala Gly	
235 240 245	
cgc gcg gcc ttg cag ctg aag ctg cgt cgg cgg ctc acg gag ctc ctg	819
Arg Ala Ala Leu Gln Leu Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu	
250 255 260 265	
ggg gcg cag gac ggg gcg ctg ctg gtg cgg ctg ctg cag gcg ctg cgc	867
Gly Ala Gln Asp Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg	
270 275 280	
gtg gcc agg atg ccc ggg ctg gag cgg agc gtc cgt gag cgc ttc ctc	915
Val Ala Arg Met Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu	
285 290 295	
cct gtg cac tgatcctggc cc	936
Pro Val His	
300	